RAW SEQUENCE LISTING   DATE: 02/08/2002
PATENT APPLICATION: US/10/040,497   TIME: 08:43:18

Input Set : N:\Crf3\RULE60\10040497.txt
Output Set: N:\CRF3\02082002\J040497.raw

ENTERED

```
        SEQUENCE LISTING
 2  (1) GENERAL INFORMATION:
 3        (i) APPLICANT: GOLD, LARRY
 4                       TUERK, CRAIG
 5       (ii) TITLE OF INVENTION: METHODS OF PRODUCING
 6                                NUCLEIC ACID LIGANDS
 7      (iii) NUMBER OF SEQUENCES: 83
 8       (iv) CORRESPONDENCE ADDRESS:
 9            (A) ADDRESSEE: Swanson & Bratschun, L.L.C.
10            (B) STREET: 8400 E. Prentice Avenue, Suite 200
11            (C) CITY: Englewood
12            (D) STATE: Colorado
13            (E) COUNTRY: USA
14            (F) ZIP: 80111
15        (v) COMPUTER READABLE FORM:
16            (A) MEDIUM TYPE: Diskette, 3 1/5 inch, 1.44 MB
17            (B) COMPUTER: IBM compatible
18            (C) OPERATING SYSTEM: MS-DOS
19            (D) SOFTWARE: WordPerfect 8.0
20       (vi) CURRENT APPLICATION DATA:
C--> 21       (A) APPLICATION NUMBER: US/10/040,497
C--> 22       (B) FILING DATE: 07-Jan-2002
C--> 39      (vii) PRIOR APPLICATION DATA:
24            (A) APPLICATION NUMBER: 09/165,616
25            (B) FILING DATE: 02-OCTOBER-1998
28            (A) APPLICATION NUMBER: 08/748,697
29            (B) FILING DATE: 13-NOVEMBER-1996
31            (A) APPLICATION NUMBER: 08/442,062
32            (B) FILING DATE: 16-MAY-1995
34            (A) APPLICATION NUMBER: 07/964,624
35            (B) FILING DATE: 21-OCTOBER-1992
37            (A) APPLICATION NUMBER: 07/714,131
38            (B) FILING DATE: 10-JUNE-1991
40            (A) APPLICATION NUMBER: 07/536,428
41            (B) FILING DATE: 11-JUNE-1990
42      (viii) ATTORNEY/AGENT INFORMATION:
43            (A) NAME: Barry J. Swanson
44            (B) REGISTRATION NUMBER: 33,215
45            (C) REFERENCE/DOCKET NUMBER: NEX05/DC-CON
46       (ix) TELECOMMUNICATION INFORMATION:
47            (A) TELEPHONE: (303) 793-3333
48            (B) TELEFAX: (303) 793-3433
49  (2) INFORMATION FOR SEQ ID NO: 1:
```

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```
50      (i) SEQUENCE CHARACTERISTICS:
51          (A) LENGTH: 21 base pairs
52          (B) TYPE: nucleic acid
53          (C) STRANDEDNESS: single
54          (D) TOPOLOGY: linear
55      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
56 UUCCGNNNNN NNNCGGGAAA A                        21
58 (2) INFORMATION FOR SEQ ID NO: 2:
59          (i) SEQUENCE CHARACTERISTICS:
60              (A) LENGTH: 24 base pairs
61              (B) TYPE: nucleic acid
62              (C) STRANDEDNESS: single
63              (D) TOPOLOGY: linear
64      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
65 UCCGNNNNNN NNCGGGAAAA NNNN                      24
67 (2) INFORMATION FOR SEQ ID NO: 3:
68          (i) SEQUENCE CHARACTERISTICS:
69              (A) LENGTH: 27 base pairs
70              (B) TYPE: nucleic acid
71              (C) STRANDEDNESS: single
72              (D) TOPOLOGY: linear
73      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
74 NNUUCCGNNN NNNNNCGGGA AAANNNN                   27
76 (2) INFORMATION FOR SEQ ID NO: 4:
77          (i) SEQUENCE CHARACTERISTICS:
78              (A) LENGTH: 57 base pairs
79              (B) TYPE: nucleic acid
80              (C) STRANDEDNESS: single
81              (D) TOPOLOGY: linear
82      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
83 GGUCCGAAGU GCAACGGGAA AAUGCACUAU GAAAGAAUUU UAUAUCUCUA   50
84 UUGAAAC                                                 57
86 (2) INFORMATION FOR SEQ ID NO: 5:
87          (i) SEQUENCE CHARACTERISTICS:
88              (A) LENGTH: 31 base pairs
89              (B) TYPE: nucleic acid
90              (C) STRANDEDNESS: single
91              (D) TOPOLOGY: linear
92      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
93 CCGGATCCGT TTCAATAGAG ATATAAAATT C             31
95 (2) INFORMATION FOR SEQ ID NO: 6:
96          (i) SEQUENCE CHARACTERISTICS:
97              (A) LENGTH: 55 base pairs
98              (B) TYPE: nucleic acid
99              (C) STRANDEDNESS: single
100             (D) TOPOLOGY: linear
101     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
102 GGGCAAGCTT TAATACGACT CACTATAGGT CCGAAGTGCA ACGGGAAAAT   50
103 GCACT                                                   55
```

```
105 (2) INFORMATION FOR SEQ ID NO: 7:
106      (i) SEQUENCE CHARACTERISTICS:
107           (A) LENGTH: 31 base pairs
108           (B) TYPE: nucleic acid
109           (C) STRANDEDNESS: single
110           (D) TOPOLOGY: linear
111      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
112 GTTTCAATAG AGATATAAAA TTCTTTCATA G                        31
114 (2) INFORMATION FOR SEQ ID NO: 8:
115      (i) SEQUENCE CHARACTERISTICS:
116           (A) LENGTH: 89 base pairs
117           (B) TYPE: nucleic acid
118           (C) STRANDEDNESS: single
119           (D) TOPOLOGY: linear
120      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
121 GTTTCAATAG AGATATAAAA TTCTTTCATA GNNNNNNNNN NNNNNNNNNN      50
122 NNNNNNNNNN NAGTGCATTT TCCCGTTGCA CTTCGGACC                 89
124 (2) INFORMATION FOR SEQ ID NO: 9:
125      (i) SEQUENCE CHARACTERISTICS:
126           (A) LENGTH: 85 base pairs
127           (B) TYPE: nucleic acid
128           (C) STRANDEDNESS: single
129           (D) TOPOLOGY: linear
130      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
131 CCCGGATCCT CTTTACCTCT GTGTGAGATA CAGAGTCCAC AAACGTGTTC      50
132 TCAATGCACC CGGTCGGAAG GCCATCAATA GTCCC                     85
134 (2) INFORMATION FOR SEQ ID NO: 10:
135      (i) SEQUENCE CHARACTERISTICS:
136           (A) LENGTH: 49 base pairs
137           (B) TYPE: nucleic acid
138           (C) STRANDEDNESS: single
139           (D) TOPOLOGY: linear
140      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
141 CCGAAGCTTA ATACGACTCA CTATAGGGAC TATTGATGGC CTTCCGACC       49
143 (2) INFORMATION FOR SEQ ID NO: 11:
144      (i) SEQUENCE CHARACTERISTICS:
145           (A) LENGTH: 25 base pairs
146           (B) TYPE: nucleic acid
147           (C) STRANDEDNESS: single
148           (D) TOPOLOGY: linear
149      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
150 CCCGGATCCT CTTTACCTCT GTGTG                                25
152 (2) INFORMATION FOR SEQ ID NO: 12:
153      (i) SEQUENCE CHARACTERISTICS:
154           (A) LENGTH: 23 base pairs
155           (B) TYPE: nucleic acid
156           (C) STRANDEDNESS: single
157           (D) TOPOLOGY: linear
158      (ix) FEATURE:
```

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PATENT APPLICATION: US/10/040,497

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```
159          (D) OTHER INFORMATION: N's at
160 positions 9-11 are base
161 paired with N's at positions 21-23
162      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:
163 UCCSANNNNN NASGGGANAA NNN                                    23
165 (2) INFORMATION FOR SEQ ID NO: 13:
166      (i) SEQUENCE CHARACTERISTICS:
167          (A) LENGTH: 66 base pairs
168          (B) TYPE: nucleic acid
169          (C) STRANDEDNESS: single
170          (D) TOPOLOGY: linear
W--> 171    (ii) MOLECULE TYPE: RNA
172    · (ix) FEATURE:
173          (D) OTHER INFORMATION: N at
174 position 29-36 can have a
175 variable length of 6-8 bases
176      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:
177 GGUCCGAAGU GCAACGGGGA AAAUGCACNN NNNNNNNNNN NNNNNNNNNN      50
178 NNNNNNNNNN NNNNNN                                           66
180 (2) INFORMATION FOR SEQ ID NO: 14:
181      (i) SEQUENCE CHARACTERISTICS:
182          (A) LENGTH: 88 base pairs
183          (B) TYPE: nucleic acid
184          (C) STRANDEDNESS: single
185          (D) TOPOLOGY: linear
186      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:
187 GGUCCGAAGU GCAACGGGAA AAUGCACUAG CUCGUGAGGC UUUCGUGCUG      50
188 UUCCGAGCUA UGAAAGAAUU UUAUAUCUCU AUUGAAAC                   88
190 (2) INFORMATION FOR SEQ ID NO: 15:
191      (i) SEQUENCE CHARACTERISTICS:
192          (A) LENGTH: 87 base pairs
193          (B) TYPE: nucleic acid
194          (C) STRANDEDNESS: single
195          (D) TOPOLOGY: linear
196      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:
197 GGUCCGAAGU GCAACGGGAA AAUGCACUGC AUGUGAGGCG GUAACGCUGU      50
198 UCCGUGCUAU GAAAGAAUUU UAUAUCUCUA UUGAAAC                    87
200 (2) INFORMATION FOR SEQ ID NO: 16:
201      (i) SEQUENCE CHARACTERISTICS:
202          (A) LENGTH: 89 base pairs
203          (B) TYPE: nucleic acid
204          (C) STRANDEDNESS: single
205          (D) TOPOLOGY: linear
206      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:
207 GGUCCGAAGU GCAACGGGAA AAUGCACUGG UGAGUGAGGC CGAUGCUGUU      50
208 CCUCGCCGCU AUGAAAGAAU UUUAUAUCUC UAUUGAAAC                  89
210 (2) INFORMATION FOR SEQ ID NO: 17:
211      (i) SEQUENCE CHARACTERISTICS:
212          (A) LENGTH: 91 base pairs
```

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/040,497

DATE: 02/08/2002
TIME: 08:43:18

Input Set : N:\Crf3\RULE60\10040497.txt
Output Set: N:\CRF3\02082002\J040497.raw

```
213           (B) TYPE: nucleic acid
214           (C) STRANDEDNESS: single
215           (D) TOPOLOGY: linear
216     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:
217  GGUCCGAAGU GCAACGGGAA AAUGCACUGA CGCGCGAGGU CUUGGUACUG        50
218  UUCCGUGGCU CUAUGAAAGA AUUUUAUAUC UCUAUUGAAA C                91
220  (2) INFORMATION FOR SEQ ID NO: 18:
221        (i) SEQUENCE CHARACTERISTICS:
222           (A) LENGTH: 89 base pairs
223           (B) TYPE: nucleic acid
224           (C) STRANDEDNESS: single
225           (D) TOPOLOGY: linear
226     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:
227  GGUCCGAAGU GCAACGGGAA AAUGCACUCU GGGUGAGACU UGAAGUCGUU        50
228  CCCCAGGUCU AUGAAAGAAU UUUAUAUCUC UAUUGAAAC                   89
230  (2) INFORMATION FOR SEQ ID NO: 19:
231        (i) SEQUENCE CHARACTERISTICS:
232           (A) LENGTH: 89 base pairs
233           (B) TYPE: nucleic acid
234           (C) STRANDEDNESS: single
235           (D) TOPOLOGY: linear
236     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:
237  GGUCCGAAGU GCAACGGGAA AAUGCACUCC CGGUGAAGCA UAAUGCUGUU        50
238  CCUGGGGUCU AUGAAAGAAU UUUAUAUCUC UAUUGAAAC                   89
240  (2) INFORMATION FOR SEQ ID NO: 20:
241        (i) SEQUENCE CHARACTERISTICS:
242           (A) LENGTH: 89 base pairs
243           (B) TYPE: nucleic acid
244           (C) STRANDEDNESS: single
245           (D) TOPOLOGY: linear
246     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:
247  GGUCCGAAGU GCAACGGGAA AAUGCACUGG GAGUGAGGUU CCCCGUUCCU        50
248  CCCGCACCCU AUGAAAGAAU UUUAUAUCUC UAUUGAAAC                   89
250  (2) INFORMATION FOR SEQ ID NO: 21:
251        (i) SEQUENCE CHARACTERISTICS:
252           (A) LENGTH: 89 base pairs
253           (B) TYPE: nucleic acid
254           (C) STRANDEDNESS: single
255           (D) TOPOLOGY: linear
256     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 21:
257  GGUCCGAAGU GCAACGGGAA AAUGCACUAG CGAUGUGAAG UGAUACUGGU        50
258  CCAUCGUGCU AUGAAAGAAU UUUAUAUCUC UAUUGAAAC                   89
260  (2) INFORMATION FOR SEQ ID NO: 22:
261        (i) SEQUENCE CHARACTERISTICS:
262           (A) LENGTH: 88 base pairs
263           (B) TYPE: nucleic acid
264           (C) STRANDEDNESS: single
265           (D) TOPOLOGY: linear
266     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 22:
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VERIFICATION SUMMARY
PATENT APPLICATION: US/10/040,497

DATE: 02/08/2002
TIME: 08:43:19

Input Set : N:\Crf3\RULE60\10040497.txt
Output Set: N:\CRF3\02082002\J040497.raw

L:21 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:22 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
L:27 M:220 C: Keyword misspelled or invalid format, [(vii) PRIOR APPLICATION DATA:]
L:171 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:171 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=13
L:412 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:412 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=36
L:426 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:426 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=37
L:437 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:437 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=38
L:459 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:459 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=39
L:909 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:909 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=83